

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/977, 015Source: 009Date Processed by STIC: 0-19-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

Or

- U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 69/977, 615	
ATTN: NEW RULES CASES	s: Please disregard english "alpha" headers. Which were inserted by Pto so	OFTWARE
1Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules Please ensure your subsequent submission is saved in ASCII text.	eras e e e e e e e e e e e e e e e e e e e
5Variable Length	Sequence(s)contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	<u></u> . i.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped	
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 Skipped Sequences (NEW RULES)	Sequence(s) missing. If Intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.	

AMC/MH - Biotechnology Systems Branch - 08/21/2001



RAW SEQUENCE LISTING

DATE: 06/19/2003 TIME: 15:21:03

PATENT APPLICATION: US/09/977,615

Input Set : A:\09977615 Sequence Listing.txt
Output Set: N:\CRF4\06192003\I977615.raw

```
3 <110> APPLICANT: Grenier, Jennifer
                                                          Does Not Comply
              Marshall, David
                                                          Corrected Diskette Needed
      5
              Prudent, James
              Richmond, Craig
      6
      7
              Roesch, Eric
      8
              Scherrer, Christopher
              Sherrill, Christopher
      9
              Ptacin, Jerod
     12 <120> TITLE OF INVENTION: SOLID SUPPORT ASSAY SYSTEMS AND METHODS UTILIZING NON-
NATURAL BASES
     14 <130> FILE REFERENCE: 031403-9012
     16 <140> CURRENT APPLICATION NUMBER: 09/977,615
     17 <141> CURRENT FILING DATE: 2001-10-15
     19 <150> PRIOR APPLICATION NUMBER: 60/240,397
     20 <151> PRIOR FILING DATE: 2000-10-14
     22 <150> PRIOR APPLICATION NUMBER: 60/282,831
     23 <151> PRIOR FILING DATE: 2001-04-10
     25 <150> PRIOR APPLICATION NUMBER: 09/861,292
     26 <151> PRIOR FILING DATE: 2001-05-18
                                                             The type of errors shown exist throughout
     28 <150> PRIOR APPLICATION NUMBER: 60/293,259
                                                             the Sequence Listing. Please check subsequent
     29 <151> PRIOR FILING DATE: 2001-05-22
                                                             sequences for similar errors.
     31 <160> NUMBER OF SEQ ID NOS: 156
     33 <170> SOFTWARE: PatentIn version 3.0
                                                       _ See Hem # 10 on Error
Summary Sheet
     35 <210> SEQ ID NO: 1
     36 <211> LENGTH: 10
     37 <212> TYPE: DNA _
     38 <213> ORGANISM: (synthetic oligonucleotide
     40 <220> FEATURE:
     41 <221> NAME/KEY: modified_base
     42 <222> LOCATION: (3)..(3)
     43 <223> OTHER INFORMATION: n represents iso-guanine
     46 <220> FEATURE:
     47 <221> NAME/KEY: modified base
     48 <222> LOCATION: (6)..(6)
     49 <223> OTHER INFORMATION: n represents iso-guanine
     52 <400> SEQUENCE: 1
                                                                                  10
W--> 53 gangtntgtc
     56 <210> SEQ ID NO: 2
     57 <211> LENGTH: 10
     58 <212> TYPE: DNA-
     59 <213> ORGANISM: synthetic oligonucleotide
     61 <220> FEATURE:
     62 <221> NAME/KEY: modified base
     63 <222> LOCATION: (2)..(2)
```

DATE: 06/19/2003

```
PATENT APPLICATION: US/09/977,615
                                                              TIME: 15:21:03
                     Input Set : A:\09977615 Sequence Listing.txt
                     Output Set: N:\CRF4\06192003\I977615.raw
     64 <223> OTHER INFORMATION: n represents iso-quanine
     67 <220> FEATURE:
     68 <221> NAME/KEY: modified base
     69 <222> LOCATION: (6)..(6)
     70 <223> OTHER INFORMATION: n represents iso-quanine
     73 <400> SEQUENCE: 2
W--> 74 cngttnttcc
                                                                                10
     77 <210> SEQ ID NO: 3
     78 <211> LENGTH: 10
     79 <212> TYPE: DNA
                                                    - See page 1
     80 <213> ORGANISM: synthetic oligonucleotide
     82 <220> FEATURE:
     83 <221> NAME/KEY: modified base
     84 <222> LOCATION: (3)..(3)
     85 <223> OTHER INFORMATION: n represents iso-guanine
     88 <220> FEATURE:
     89 <221> NAME/KEY: modified_base
     90 <222> LOCATION: (7)..(7)
     91 <223> OTHER INFORMATION: n represents iso-guanine
     94 <400> SEQUENCE: 3
W--> 95 ggnttgntag
                                                                                10
     98 <210> SEQ ID NO: 4
     99 <211> LENGTH: 10
     100 <212> TYPE: DNA
     101 <213> ORGANISM: synthetic oligonucleotide
     103 <220> FEATURE:
     104 <221> NAME/KEY: modified base
     105 <222> LOCATION: (4)..(4)
     106 <223> OTHER INFORMATION: n represents iso-quanine
     109 <220> FEATURE:
     110 <221> NAME/KEY: modified base
     111 <222> LOCATION: (6)..(6)
     112 <223> OTHER INFORMATION: n represents iso-guanine
     115 <400> SEQUENCE: 4
W--> 116 cttngntctc
                                                                                10
     119 <210> SEQ ID NO: 5
     120 <211> LENGTH: 10
     121 <212> TYPE: DNA
     122 <213> ORGANISM: (synthetic oligonucleotide
     124 <220> FEATURE:
     125 <221> NAME/KEY: modified base
     126 <222> LOCATION: (2)..(2)
     127 <223> OTHER INFORMATION: n represents iso-guanine
     130 <220> FEATURE:
     131 <221> NAME/KEY: modified base
     132 <222> LOCATION: (6)..(6)
     133 <223> OTHER INFORMATION: n represents iso-quanine
     136 <400> SEQUENCE: 5
W--> 137 cntcangaac
                                                                                10
```

RAW SEQUENCE LISTING

DATE: 06/19/2003

TIME: 15:21:03

Input Set: A:\09977615 Sequence Listing.txt Output Set: N:\CRF4\06192003\I977615.raw 140 <210> SEQ ID NO: 6 141 <211> LENGTH: 10 142 <212> TYPE: DNA 143 <213> ORGANISM: (synthetic oligonucleotide 145 <220> FEATURE: 146 <221> NAME/KEY: modified base 147 <222> LOCATION: (5)..(5) 148 <223> OTHER INFORMATION: n represents iso-quanine 151 <220> FEATURE: 152 <221> NAME/KEY: modified base 153 <222> LOCATION: (8)..(8) 154 <223> OTHER INFORMATION: n represents iso-quanine 157 <400> SEQUENCE: 6 W--> 158 gtagntangc 10 161 <210> SEQ ID NO: 7 162 <211> LENGTH: 10 163 <212> TYPE: DNA -- see page 164 <213> ORGANISM: (synthetic oligonucleotide) 166 <220> FEATURE: 167 <221> NAME/KEY: modified_base 168 <222> LOCATION: (4)..(4) 169 <223> OTHER INFORMATION: n represents iso-guanine 172 <220> FEATURE: 173 <221> NAME/KEY: modified base 174 <222> LOCATION: (6)..(6) 175 <223> OTHER INFORMATION: n represents iso-quanine 178 <400> SEQUENCE: 7 W--> 179 ggangntaac 10 182 <210> SEQ ID NO: 8 183 <211> LENGTH: 10 184 <212> TYPE: DNA_ 185 <213> ORGANISM: (synthetic oligonucleotide 187 <220> FEATURE: 188 <221> NAME/KEY: modified base 189 <222> LOCATION: (2)..(2) 190 <223> OTHER INFORMATION: n represents iso-quanine 193 <220> FEATURE: 194 <221> NAME/KEY: modified base 195 <222> LOCATION: (7)...(7) 196 <223> OTHER INFORMATION: n represents iso-quanine 199 <400> SEQUENCE: 8 W--> 200 cngtatngtg 10 203 <210> SEQ ID NO: 9 204 <211> LENGTH: 10 205 <212> TYPE: DNA 206 <213> ORGANISM: /synthetic oligonucleotide 208 <220> FEATURE: 209 <221> NAME/KEY: modified_base 210 <222> LOCATION: (4)..(4)

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/977,615

DATE: 06/19/2003

TIME: 15:21:03

Input Set : A:\09977615 Sequence Listing.txt Output Set: N:\CRF4\06192003\I977615.raw 211 <223> OTHER INFORMATION: n represents iso-guanine 214 <220> FEATURE: 215 <221> NAME/KEY: modified base 216 <222> LOCATION: (10)..(10) 217 <223> OTHER INFORMATION: n represents iso-quanine 220 <400> SEQUENCE: 9 W--> 221 catnggtang 10 224 <210> SEQ ID NO: 10 225 <211> LENGTH: 10 226 <212> TYPE: DNA 227 <213> ORGANISM: synthetic oligonucleotide 229 <220> FEATURE: 230 <221> NAME/KEY: modified base 231 <222> LOCATION: (5)..(5) 232 <223> OTHER INFORMATION: n represents iso-guanine 235 <220> FEATURE: 236 <221> NAME/KEY: modified_base 237 <222> LOCATION: (9)..(9) 238 <223> OTHER INFORMATION: n represents iso-guanine 241 <400> SEQUENCE: 10 W--> 242 gattntcgnc 10 245 <210> SEQ ID NO: 11 246 <211> LENGTH: 10 - Su page 1 247 <212> TYPE: DNA 248 <213> ORGANISM: synthetic oligonucleotide 250 <220> FEATURE: 251 <221> NAME/KEY: modified base 252 <222> LOCATION: (4)..(4) 253 <223> OTHER INFORMATION: n represents iso-guanine 256 <220> FEATURE: 257 <221> NAME/KEY: modified base 258 <222> LOCATION: (6)..(6) 259 <223> OTHER INFORMATION: n represents iso-quanine 262 <400> SEQUENCE: 11 W--> 263 gttnangacc 10 266 <210> SEO ID NO: 12 267 <211> LENGTH: 10 268 <212> TYPE: DNA 269 <213> ORGANISM: (synthetic oligonucleotide) 271 <220> FEATURE: 272 <221> NAME/KEY: modified_base 273 <222> LOCATION: (2)..(2) 274 <223> OTHER INFORMATION: n represents iso-guanine 277 <220> FEATURE: 278 <221> NAME/KEY: modified base 279 <222> LOCATION: (6)..(6) 280 <223> OTHER INFORMATION: n represents iso-quanine 283 <400> SEQUENCE: 12 W--> 284 cngaangatc 10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/977,615

RAW SEQUENCE LISTING DATE: 06/19/2003 PATENT APPLICATION: US/09/977,615 TIME: 15:21:03

Input Set : A:\09977615 Sequence Listing.txt
Output Set: N:\CRF4\06192003\I977615.raw

```
287 <210> SEQ ID NO: 13
     288 <211> LENGTH: 10
     289 <212> TYPE: DNA
     290 <213> ORGANISM:/synthetic oligonucleotide
     292 <220> FEATURE:
     293 <221> NAME/KEY: modified base
     294 <222> LOCATION: (4)..(4)
     295 <223> OTHER INFORMATION: n represents iso-quanine
     298 <220> FEATURE:
     299 <221> NAME/KEY: modified base
     300 <222> LOCATION: (9)..(9)
     301 <223> OTHER INFORMATION: n represents iso-guanine
     304 <400> SEQUENCE: 13
W--> 305 caantacgnc
                                                                                10
     308 <210> SEQ ID NO: 14
     309 <211> LENGTH: 10
     310 <212> TYPE: DNA, ___
     311 <213> ORGANISM: synthetic oligonucleotide See page (
     313 <220> FEATURE:
     314 <221> NAME/KEY: modified base
     315 <222> LOCATION: (4)..(4)
     316 <223> OTHER INFORMATION: n represents iso-guanine
     319 <220> FEATURE:
     320 <221> NAME/KEY: modified base
     321 <222> LOCATION: (8)..(8)
     322 <223> OTHER INFORMATION: n represents iso-quanine
     325 <400> SEQUENCE: 14
W--> 326 cggnatanac
                                                                                10
     329 <210> SEQ ID NO: 15
     330 <211> LENGTH: 10
     331 <212> TYPE: DNA
     332 <213> ORGANISM: synthetic oligonucleotide
     334 <220> FEATURE:
     335 <221> NAME/KEY: modified base
     336 <222> LOCATION: (2)..(2)
     337 <223> OTHER INFORMATION: n represents iso-quanine
     340 <220> FEATURE:
     341 <221> NAME/KEY: modified base
     342 <222> LOCATION: (6)..(6)
     343 <223> OTHER INFORMATION: n represents iso-guanine
     346 <220> FEATURE:
     347 <221> NAME/KEY: modified_base
     348 <222> LOCATION: (7)..(7)
     349 <223> OTHER INFORMATION: n represents iso-quanine
     352 <400> SEQUENCE: 15
W--> 353 gnaaannagg
                                                                                10
     356 <210> SEQ ID NO: 16
     357 <211> LENGTH: 10
     358 <212> TYPE: DNA
```

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/19/2003 PATENT APPLICATION: US/09/977,615 TIME: 15:21:04

Input Set : A:\09977615 Sequence Listing.txt
Output Set: N:\CRF4\06192003\I977615.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:1; N Pos. 3,6
Seq#:2; N Pos. 2,6
Seq#:3; N Pos. 3,7
Seq#:4; N Pos. 4,6
Seq#:5; N Pos. 2,6
Seq#:6; N Pos. 5,8
Seq#:7; N Pos. 4,6
Seq#:8; N Pos. 2,7
Seq#:9; N Pos. 4,9
Seq#:10; N Pos. 5,9
Seq#:11; N Pos. 4,6
Seq#:12; N Pos. 2,6
Seq#:13; N Pos. 4,9
Seq#:14; N Pos. 4,8
Seq#:15; N Pos. 2,6,7
Seq#:16; N Pos. 4,8,9
Seq#:17; N Pos. 2,6,9
Seq#:18; N Pos. 3,6
Seq#:19; N Pos. 3,4,7
Seq#:20; N Pos. 3,6
Seq#:21; N Pos. 5,8,9
Seq#:22; N Pos. 5,7
Seq#:23; N Pos. 5,8
Seq#:24; N Pos. 5,8
Seq#:25; N Pos. 5,7,8
Seq#:26; N Pos. 2,7
Seq#:27; N Pos. 2,4,9
Seq#:28; N Pos. 2,3,7
Seq#:29; N Pos. 5,8
Seq#:30; N Pos. 5,7
Seq#:31; N Pos. 3,7
Seq#:32; N Pos. 4,8
Seq#:33; N Pos. 4,9
Seq#:34; N Pos. 4,7,9
Seq#:35; N Pos. 2,7
Seq#:36; N Pos. 3,7
Seq#:37; N Pos. 2,7
Seq#:38; N Pos. 4,9
Seq#:39; N Pos. 3,4,9
Seq#:40; N Pos. 2,6
Seq#:41; N Pos. 2,6,9
Seq#:42; N Pos. 3,6
Seq#:43; N Pos. 4,8
Seq#:44; N Pos. 5,8
```

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/19/2003 PATENT APPLICATION: US/09/977,615 TIME: 15:21:04

Input Set : A:\09977615 Sequence Listing.txt
Output Set: N:\CRF4\06192003\I977615.raw

Seq#:45; N Pos. 5,7 Seq#:46; N Pos. 4,8 Seq#:47; N Pos. 2,7 Seq#:48; N Pos. 3,6 Seq#:49; N Pos. 3,7 Seq#:50; N Pos. 3,4,9 Seq#:51; N Pos. 2,7

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/977,615 TIME: 15:21:04

DATE: 06/19/2003

Input Set : A:\09977615 Sequence Listing.txt
Output Set: N:\CRF4\06192003\I977615.raw

```
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:158 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:242 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:263 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:284 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:407 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:455 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:503 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:503 M:341 W: (40) "" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:524 M:341 W: (46) "" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:545 M:341 W: (46) "" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:566 M:341 W: (46) "" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:593 M:341 W: (46) "" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:614 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:668 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:710 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:752 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
L:800 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:821 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:842 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:863 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0
L:884 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0
L:911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0
L:932 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:0
L:959 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
L:980 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:0
L:1001 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0
L:1022 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0
L:1043 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:0
L:1064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 after pos.:0
L:1085 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:0
L:1106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:0
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/977,615

DATE: 06/19/2003 TIME: 15:21:04

Input Set : A:\09977615 Sequence Listing.txt
Output Set: N:\CRF4\06192003\I977615.raw

L:1127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:0 L:1154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0